



SEQUENCE LISTING

<110> Berchtold, Peter  
Escher, Robert F. A.

<120> ANTI-GPIIB/IIIA RECOMBINANT ANTIBODIES

<130> 100564-09049

<140> 09/424,840  
<141> 1999-12-03

<150> DE 19820663.1  
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<150> DE 19755227.7  
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<151> 1997-06-06

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<170> PatentIn version 3.1

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Thr Leu Ser Leu Asn Cys Thr Val Ser Gly Arg Ser Ile Ser Gly Tyr		
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tct tgg aga tgg atc cgg cag tct cca ggg aag gga cta gag tgg att	144	
Ser Trp Arg Trp Ile Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Ile		
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Gly Asp Ile Ser Tyr Ser Gly Ser Thr Lys Tyr Lys Pro Ser Leu Arg		
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agt cga gtc acc ctg tca gta gac acg tcc aag aac cag ttc tcc ctg	240	
Ser Arg Val Thr Leu Ser Val Asp Thr Ser Lys Asn Gln Phe Ser Leu		
65 70 75 80		
aag ctg aat tcg gtg acc gct gcg gac acg gcc gtc tat tac tgt gcg	288	
Lys Leu Asn Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala		
85 90 95		
cga gtc ttg ccc ttt gac ccg atc tcg atg gac gtc tgg ggc aaa ggg	336	
Arg Val Leu Pro Phe Asp Pro Ile Ser Met Asp Val Trp Gly Lys Gly		
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ser Trp Arg Trp Ile Arg Gln Ser Pro Gly Lys Gly Leu Glu Trp Ile		
35 40 45		

Gly Asp Ile Ser Tyr Ser Gly Ser Thr Lys Tyr Lys Pro Ser Leu Arg  
50 55 60

Ser Arg Val Thr Leu Ser Val Asp Thr Ser Lys Asn Gln Phe Ser Leu  
65 70 75 80

Lys Leu Asn Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala  
85 90 95

Arg Val Leu Pro Phe Asp Pro Ile Ser Met Asp Val Trp Gly Lys Gly  
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Thr Thr Val Thr Val Ser Ser  
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acc atc tct tgt tct ggg agc agc tcc aac atc aga agt aat cct gtt  
Thr Ile Ser Cys Ser Gly Ser Ser Asn Ile Arg Ser Asn Pro Val  
20 25 30

agc tgg tat cac cag gtc cca ggc acg gcc ccc aaa ctc ctc atc ttt  
Ser Trp Tyr His Gln Val Pro Gly Thr Ala Pro Lys Leu Leu Ile Phe  
35 40 45

ggt agt cat cag cgg ccc tca ggg gtc cct gac cga ttc tct ggc tcc  
Gly Ser His Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60

aag tcg ggc acc tcc gcc tcc ctg gcc atc cgt ggg ctc caa tct ggg  
Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Arg Gly Leu Gln Ser Gly  
65 70 75 80

gat gct ggt gac tat tac tgt gca aca tgg gat gac ggc ctc aat ggt  
Asp Ala Gly Asp Tyr Tyr Cys Ala Thr Trp Asp Asp Gly Leu Asn Gly  
85 90 95

48

96

144

192

240

288

333

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<213> Homo sapiens

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20 25 30

Ser Trp Tyr His Gln Val Pro Gly Thr Ala Pro Lys Leu Leu Ile Phe  
35 40 45

Gly Ser His Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60

Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Arg Gly Leu Gln Ser Gly  
65 70 75 80

Asp Ala Gly Asp Tyr Tyr Cys Ala Thr Trp Asp Asp Gly Leu Asn Gly  
85 90 95

Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Ser Gln Pro  
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr		
20 25 30		
gct atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg gag tgg gtg	144	
Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val		
35 40 45		
gca gtt ata tca tat gat gga agc aat aaa tac tac gca gac tcc gta	192	
Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val		
50 55 60		
aag ggc cga ttc gcc atc tcc aga gac aat tcc aag aac acg ctg tat	240	
Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr		
65 70 75 80		
ctg caa atg aac agc ctg aga gct gag gac acg gct gta tat tac tgt	288	
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys		
85 90 95		
gcg aga gcg ctg ggg agc tgg ggg ggt tgg gac cac tac atg gac gtc	336	
Ala Arg Ala Leu Gly Ser Trp Gly Gly Trp Asp His Tyr Met Asp Val		
100 105 110		
tgg ggc aaa ggg acc acg gtc acc gtc tcc tca	369	
Trp Gly Lys Gly Thr Thr Val Thr Val Ser Ser		
115 120		

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20 25 30						
Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val						
35 40 45						
Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val						
50 55 60						
Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr						
5						

65

70

75

80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Arg Ala Leu Gly Ser Trp Gly Gly Trp Asp His Tyr Met Asp Val  
100 105 110

Trp Gly Lys Gly Thr Thr Val Thr Val Ser Ser  
115 120

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acc atc tct tgt tct gga agc agc tcc aac atc gga agt aat act gta 96  
Thr Ile Ser Cys Ser Gly Ser Ser Asn Ile Gly Ser Asn Thr Val  
20 25 30  
aac tgg tac cag cag ctc cca gga acg gcc ccc aaa ctc ctc atc tat 144  
Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr  
35 40 45  
agt aat aat cag cgg ccc tca ggg gtc cct gac cga ttc tct ggc tcc 192  
Ser Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60  
aag tct ggc acc tca gcc tcc ctg gcc atc agt ggg ctc cag tct gag 240  
Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln Ser Glu  
65 70 75 80  
gat gag gct gat tat tac tgt gca gca tgg gat gac agc ctg aat ggt 288  
Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu Asn Gly  
85 90 95  
tgg gtg ttc ggc gga ggg acc aag ctg acc gtc cta ggt cag ccc 333  
Trp Val Phe Gly Gly Thr Lys Leu Thr Val Leu Gly Gln Pro  
100 105 110

<210> 8  
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<400> 8  
val val Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln Arg Val  
1 5 10 15  
Thr Ile Ser Cys Ser Gly Ser Ser Asn Ile Gly Ser Asn Thr Val  
20 25 30  
Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr  
35 40 45  
Ser Asn Asn Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60  
Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln Ser Glu  
65 70 75 80  
Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu Asn Gly  
85 90 95  
Trp Val Phe Gly Gly Thr Lys Leu Thr Val Leu Gly Gln Pro  
100 105 110

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Gln Val Lys Leu Leu Glu Ser Gly Gly Leu Val His Pro Gly Gly  
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tcc ctg aga ctc tct tgt gca gcc tct gga ttt acg ttt gac aac ttt				96
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Asn Phe				
20 25 30				
gcc atg agc tgg gtc cgc cag gct cca ggg aag ggg ctg gag tgg gtc				144
Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val				
35 40 45				
tca ggc att agt ggt ggt ctt ttg aca cac tac gca gac tcc gtc				192
Ser Gly Ile Ser Gly Gly Leu Leu Thr His Tyr Ala Asp Ser Val				
50 55 60				
aag ggc cgg ttc acc atc tcc aga aac aat tcc agg aac act gta tac				240
Lys Gly Arg Phe Thr Ile Ser Arg Asn Asn Ser Arg Asn Thr Val Tyr				
65 70 75 80				
cta caa atg aac agc ctg aga gcc gaa gac acg gcc gtg tat tat tgt				288
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys				
85 90 95				
gtg aga gat ctg ggc tat aga gta ctt tcg act ttt act ttt gat atc				336
Val Arg Asp Leu Gly Tyr Arg Val Leu Ser Thr Phe Thr Phe Asp Ile				
100 105 110				
tgg ggc cag ggg aca aag gtc acc gtc tct tca				369
Trp Gly Gln Gly Thr Lys Val Thr Val Ser Ser				
115 120				

<210> 10  
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20 25 30				
Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val				
35 40 45				
Ser Gly Ile Ser Gly Gly Leu Leu Thr His Tyr Ala Asp Ser Val				
50 55 60				
Lys Gly Arg Phe Thr Ile Ser Arg Asn Asn Ser Arg Asn Thr Val Tyr				
65 70 75 80				
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys				
8				

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100	105	110
Trp Gly Gln Gly Thr Lys Val Thr Val Ser Ser		
115	120	

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 <212> DNA  
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115

120

125

<210> 12

<211> 125

<212> PRT

<213> Homo sapiens

<400> 12

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20 25 30

val Pro Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu Met Ile  
35 40 45

Tyr Glu Gly Ser Lys Arg Pro Ser Gly val Ser Asn Arg Phe Ser Gly  
50 55 60

Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu Gln Ala  
65 70 75 80

Glu Asp Glu Ala Glu Tyr Tyr Cys Cys Ser Tyr Val His Ser Ser Thr  
85 90 95

Asn Trp Val Phe Gly Gly Thr Lys Leu Thr Val Leu Gly Gln Pro  
100 105 110

Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser  
115 120 125

<210> 13

<211> 366

<212> DNA

<213> Homo sapiens

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<221> CDS

<222> (1)..(366)

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acc ctg tct ctc acc tgc act gtc tct gat gtc tcc atc aga agt cat	96
Thr Leu Ser Leu Thr Cys Thr Val Ser Asp Val Ser Ile Arg Ser His	
20 25 30	
tac tgg agt tgg ctc cgg cag ccc cca ggg aag gga ctg gag tgg att	144
Tyr Trp Ser Trp Leu Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Ile	
35 40 45	
ggg ttt atc tat gac ggt gcg aga acc agg ttc aac ccc tcc ctc agg	192
Gly Phe Ile Tyr Asp Gly Ala Arg Thr Arg Phe Asn Pro Ser Leu Arg	
50 55 60	
agt cga gtc tcc ctt tca atg gac cca tcc aag aag cag ttt tcc ctg	240
Ser Arg Val Ser Leu Ser Met Asp Pro Ser Lys Lys Gln Phe Ser Leu	
65 70 75 80	
aaa ctg ggg tct gtg acc gct gcg gac tcg gcc gtc tac tac tgt gcg	288
Lys Leu Gly Ser Val Thr Ala Ala Asp Ser Ala Val Tyr Tyr Cys Ala	
85 90 95	
aga gac gcg gat gga gat ggc ttc agc cca tac tac ttt ccc tac tgg	336
Arg Asp Ala Asp Gly Asp Gly Phe Ser Pro Tyr Tyr Phe Pro Tyr Trp	
100 105 110	
ggc cag gga atc ccg gtc tcc gtc tcc tcg	366
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115 120	

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Tyr Trp Ser Trp Leu Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Ile	
35 40 45	
Gly Phe Ile Tyr Asp Gly Ala Arg Thr Arg Phe Asn Pro Ser Leu Arg	
50 55 60	

Ser Arg Val Ser Leu Ser Met Asp Pro Ser Lys Lys Gln Phe Ser Leu  
65 70 75 80

Lys Leu Gly Ser Val Thr Ala Ala Asp Ser Ala Val Tyr Tyr Cys Ala  
85 90 95

Arg Asp Ala Asp Gly Asp Gly Phe Ser Pro Tyr Tyr Phe Pro Tyr Trp  
100 105 110

Gly Gln Gly Ile Pro Val Ser Val Ser Ser  
115 120

<210> 15

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<212> DNA

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tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt agc tat	96
Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr	
20 25 30	
act atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg gag tgg gtg	144
Thr Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val	
35 40 45	
gca ctt ata tca tat gat gga agc aat aaa tac tac gca gac tcc gtg	192
Ala Leu Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val	
50 55 60	
aag ggc cga ttc gcc atc tcc aga gac aat tcc aag aac acg cta tat	240
Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr	
65 70 75 80	
ctg caa atg aac agc ctg aga gct gag gac acg gct gtg tat tac tgt	288
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys	
85 90 95	
gcg aaa gat ggc cgg agt ggg agc tac gcc agg ttc gac ggt atg gac	336
12	

Ala Lys Asp Gly Arg Ser Gly Ser Tyr Ala Arg Phe Asp Gly Met Asp  
100 105 110

372

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20 25 30

Thr Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ala Leu Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Ala Lys Asp Gly Arg Ser Gly Ser Tyr Ala Arg Phe Asp Gly Met Asp  
100 105 110

Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser  
115 120

<210> 17

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<212> PRT

<213> Homo sapiens

<400> 18

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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Asp Tyr  
20 25 30

Ala Leu His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
14

35

40

45

Ser Gly Ile Ser Trp Asp Ser Gly Thr Ile Gly Tyr Ala Asp Ser Val  
 50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr  
 65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Leu Tyr Tyr Cys  
 85 90 95

Val Lys Asp Met Gly Ser Ser Val Val Ala Thr Tyr Asn Ala Phe Asp  
 100 105 110

Ile Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser  
 115 120

&lt;210&gt; 19

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&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;220&gt;

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&lt;223&gt;

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1 5 10 15	
acc ctg tcc ctc acc tgc act gtc tct ggt ggc tcc ttc agt act tac	96
Thr Leu Ser Leu Thr Cys Thr Val Ser Gly Gly Ser Phe Ser Thr Tyr	
20 25 30	
tat tgg agc tgg atc cgg cag ccc cca ggg aag gga ctg gag tgg att	144
Tyr Trp Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Ile	
35 40 45	
ggg tat atc tat tac agt ggg aac acc aac tac aac ccc tcc ctc aag	192
Gly Tyr Ile Tyr Tyr Ser Gly Asn Thr Asn Tyr Asn Pro Ser Leu Lys	
50 55 60	
agt cga gcc acc ata tca gta gac acg tcc aag aac cag ttc tcc ctg	240
Ser Arg Ala Thr Ile Ser Val Asp Thr Ser Lys Asn Gln Phe Ser Leu	
65 70 75 80	
15	

aag ctg agc tct gtt acc gcc gca gac acg gcc gta tat tac tgt gcg	288
Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala	
85 90 95	
aga ctg cgt aac gat ggc tgg aat gat ggc ttt gat atc tgg ggc caa	336
Arg Leu Arg Asn Asp Gly Trp Asn Asp Gly Phe Asp Ile Trp Gly Gln	
100 105 110	
ggg aca atg gtc acc gtc tct tca	360
Gly Thr Met Val Thr Val Ser Ser	
115 120	

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<400> 20	
Gln Val Lys Leu Leu Glu Ser Gly Pro Gly Leu Val Lys Pro Ser Glu	
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20 25 30	
Tyr Trp Ser Trp Ile Arg Gln Pro Pro Gly Lys Gly Leu Glu Trp Ile	
35 40 45	
Gly Tyr Ile Tyr Tyr Ser Gly Asn Thr Asn Tyr Asn Pro Ser Leu Lys	
50 55 60	
Ser Arg Ala Thr Ile Ser Val Asp Thr Ser Lys Asn Gln Phe Ser Leu	
65 70 75 80	
Lys Leu Ser Ser Val Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala	
85 90 95	
Arg Leu Arg Asn Asp Gly Trp Asn Asp Gly Phe Asp Ile Trp Gly Gln	
100 105 110	
Gly Thr Met Val Thr Val Ser Ser	
115 120	

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	1	5 10 15	
	tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc agt gac tat	96	
	Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asp Tyr		
	20 25 30		
	ggc atg cac tgg gtc cgc cag gct cca ggc aag ggg ctg gag tgg gtg	144	
	Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val		
	35 40 45		
	gca gct ata tca tat gat gga agt aac aaa tac tat gca gac tcc gtg	192	
	Ala Ala Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val		
	50 55 60		
	aag ggc cga ttc tcc atc tcc aga gac aat tcc aac aat acg cta tat	240	
	Lys Gly Arg Phe Ser Ile Ser Arg Asp Asn Ser Asn Asn Thr Leu Tyr		
	65 70 75 80		
	ctg caa atg agc acc ctg aga gct gag gac acg gct gtc tat ttc tgt	288	
	Leu Gln Met Ser Thr Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys		
	85 90 95		
	gcg aga gat tcg gaa acg gca ata gcg gca gct gga cgg ttt gat atc	336	
	Ala Arg Asp Ser Glu Thr Ala Ile Ala Ala Ala Gly Arg Phe Asp Ile		
	100 105 110		
	tgg ggc caa ggg aca atg gtc acc gtc tct tca	369	
	Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser		
	115 120		

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<400> 22

Gln Val Lys Leu Leu Glu Ser Gly Gly Val Val Gln Pro Gly Arg		
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asp Tyr  
 20 25 30  
 Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
 35 40 45  
 Ala Ala Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val  
 50 55 60  
 Lys Gly Arg Phe Ser Ile Ser Arg Asp Asn Ser Asn Asn Thr Leu Tyr  
 65 70 75 80  
 Leu Gln Met Ser Thr Leu Arg Ala Glu Asp Thr Ala Val Tyr Phe Cys  
 85 90 95  
 Ala Arg Asp Ser Glu Thr Ala Ile Ala Ala Ala Gly Arg Phe Asp Ile  
 100 105 110  
 Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser  
 115 120

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1 5 10 15	
Gln Val Lys Leu Leu Glu Ser Gly Ala Glu Val Lys Lys Pro Gly Ser	
tcg gtg atg gtc tcc tgc aag gct tct gga ggc acc ttc agc agc cat	96
20 25 30	
Ser Val Met Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser His	
act atc agc tgg gtg cgg cag gcc cct gga caa ggc ctt gag tgg atg	144
35 40 45	
Thr Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met	

gga ggg atc acc cct atc ttt ggt aca gtc aac tac gca cag aag ttc Gly Gly Ile Thr Pro Ile Phe Gly Thr Val Asn Tyr Ala Gln Lys Phe 50 55 60	192
cag ggc aga gtc acc att acc gcg gac gaa ccc acg agc aca gcc tac Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Pro Thr Ser Thr Ala Tyr 65 70 75 80	240
atg gaa ctg agg agc ctg aca tct gac gac tcg ggc atc tat tac tgt Met Glu Leu Arg Ser Leu Thr Ser Asp Asp Ser Gly Ile Tyr Tyr Cys 85 90 95	288
gcg aga gaa gat ggc act aca gta cca agt caa ccc ctt gag ttc tgg Ala Arg Glu Asp Gly Thr Thr Val Pro Ser Gln Pro Leu Glu Phe Trp 100 105 110	336
ggc cag gga acc cgg gtc acc gtc tcc tct Gly Gln Gly Thr Arg Val Thr Val Ser Ser 115 120	366

<210> 24  
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<400> 24

Gln Val Lys Leu Leu Glu Ser Gly Ala Glu Val Lys Lys Pro Gly Ser 1 5 10 15
Ser Val Met Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Ser His 20 25 30
Thr Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met 35 40 45
Gly Gly Ile Thr Pro Ile Phe Gly Thr Val Asn Tyr Ala Gln Lys Phe 50 55 60
Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Pro Thr Ser Thr Ala Tyr 65 70 75 80
Met Glu Leu Arg Ser Leu Thr Ser Asp Asp Ser Gly Ile Tyr Tyr Cys 85 90 95
Ala Arg Glu Asp Gly Thr Thr Val Pro Ser Gln Pro Leu Glu Phe Trp 100 105 110
Gly Gln Gly Thr Arg Val Thr Val Ser Ser 115 120

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tcc ctg aga ctc tcc tgt tca gcc tct gga ttc acc ttc aat aaa tat Ser Leu Arg Leu Ser Cys Ser Ala Ser Gly Phe Thr Phe Asn Lys Tyr 20 25 30	96
gca ata cac tgg gtc cgc cag gct cca ggg aag gga ctg gaa tat gtt Ala Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val 35 40 45	144
tca gct att agt agt aat ggg ggt aac aca tac tac gca gac tcc gtg Ser Ala Ile Ser Ser Asn Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val 50 55 60	192
aag ggc aga ttc acc atc tcc aga gac aat tcc aag aac acg gtc tat Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Val Tyr 65 70 75 80	240
ctt caa atg agc agt ctg aga gct gag gac acg gct gtc tat tac tgt Leu Gln Met Ser Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95	288
gtt aga gga agt ggg agc tac tta gga tac tac ttt gac tac tgg ggc Val Arg Gly Ser Gly Ser Tyr Leu Gly Tyr Tyr Phe Asp Tyr Trp Gly 100 105 110	336
cag gga acc ctg gtc acc gtc tcc tca Gln Gly Thr Leu Val Thr Val Ser Ser	363
115 120	

<210> 26  
 <211> 121  
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 <213> Homo sapiens

<400> 26

Gln Val Lys Leu Leu Glu Ser Gly Gly Leu Val Gln Pro Gly Gly  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ser Ala Ser Gly Phe Thr Phe Asn Lys Tyr  
20 25 30

Ala Ile His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val  
35 40 45

Ser Ala Ile Ser Ser Asn Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Val Tyr  
65 70 75 80

Leu Gln Met Ser Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Val Arg Gly Ser Gly Ser Tyr Leu Gly Tyr Tyr Phe Asp Tyr Trp Gly  
100 105 110

Gln Gly Thr Leu Val Thr Val Ser Ser  
115 120

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<211> 366

<212> DNA

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<222> (1)..(366)

<223>

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1 5 10 15

acg att acc tgt ggg gga tac aag att gga agt aaa agt gtc cac tgg  
21

48

96

Thr Ile Thr Cys Gly Gly Tyr Lys Ile Gly Ser Lys Ser Val His Trp	25	30	
tac caa cag aag cca ggc cag gcc cct gta ttg gtc gtc tat gag gat	35	40	144
Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Val Tyr Glu Asp	45		
tcc tac cgg ccc tca gag atc cct gag cga ttc tct ggc tcc aac tct	50	55	192
Ser Tyr Arg Pro Ser Glu Ile Pro Glu Arg Phe Ser Gly Ser Asn Ser	60		
ggg aac atg gcc acc ctg acc atc acc ggg gtc gaa gcc ggg gat gag	65	70	240
Gly Asn Met Ala Thr Leu Thr Ile Thr Gly Val Glu Ala Gly Asp Glu	75	80	
gcc gac tac tac tgt cag gtg tgg gat aat act aat gat cag acg ata	85	90	288
Ala Asp Tyr Tyr Cys Gln Val Trp Asp Asn Thr Asn Asp Gln Thr Ile	95		
ttc ggc gga ggg acc aag ctg acc gtc cta cgt cag ccc aag gct gcc	100	105	336
Phe Gly Gly Thr Lys Leu Thr Val Leu Arg Gln Pro Lys Ala Ala	110		
ccc tcg gtc act ctg ttc ccg ccc tcc tct	115	120	366
Pro Ser Val Thr Leu Phe Pro Pro Ser Ser			

<210> 28

<211> 122

<212> PRT

<213> Homo sapiens

<400> 28

val val Thr Gln Pro Pro Ser Val Ser Val Ala Pro Arg Gln Thr Ala	15		
1	5	10	

Thr Ile Thr Cys Gly Gly Tyr Lys Ile Gly Ser Lys Ser Val His Trp	25	30	
20			

Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Val Tyr Glu Asp	35	40	
35		45	

Ser Tyr Arg Pro Ser Glu Ile Pro Glu Arg Phe Ser Gly Ser Asn Ser	55	60	
50			

Gly Asn Met Ala Thr Leu Thr Ile Thr Gly Val Glu Ala Gly Asp Glu	70	75	
65		80	

Ala Asp Tyr Tyr Cys Gln Val Trp Asp Asn Thr Asn Asp Gln Thr Ile	85	90	
		95	

Phe Gly Gly Thr Lys Leu Thr Val Leu Arg Gln Pro Lys Ala Ala  
100 105 110

Pro Ser Val Thr Leu Phe Pro Pro Ser Ser  
115 120

<210> 29

<211> 366

<212> DNA

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<223>

<400> 29	48
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Gln Val Lys Leu Leu Glu Ser Gly Ala Glu Val Lys Lys Pro Gly Ala	
1 5 10 15	
tca gtg aag gtc tcc tgc aag gtt tcc gga tac acc ctc act gaa tta	96
Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu	
20 25 30	
tcc atg cac tgg gtg cga cag gct cct gga aaa ggg ctt gag tgg atg	144
Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Leu Glu Trp Met	
35 40 45	
gga ggt ttt gat cct gaa gat ggt gaa aca atc tac gca cag aaa ttc	192
Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe	
50 55 60	
cag ggc aga gtc acc atg acc gag gac aca tct aca gac acg gcc tac	240
Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr	
65 70 75 80	
atg gag ctg agc agc ctg aga tct gag gac acg gcc gtg tat tac tgt	288
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys	
85 90 95	
gag aca ggt ctg agg tcg tac aac tat ggt cgt aac ctt gac tat tgg	336
Glu Thr Gly Leu Arg Ser Tyr Asn Tyr Gly Arg Asn Leu Asp Tyr Trp	
100 105 110	
ggc cag gga acc ctg gtc acc gtc tcc tca	366
Gly Gln Gly Thr Leu Val Thr Val Ser Ser	
115 120	

<210> 30

<211> 122

<212> PRT

<213> Homo sapiens

<400> 30

Gln Val Lys Leu Leu Glu Ser Gly Ala Glu Val Lys Lys Pro Gly Ala  
1 5 10 15

Ser Val Lys Val Ser Cys Lys Val Ser Gly Tyr Thr Leu Thr Glu Leu  
20 25 30 35

Ser Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Met  
35 40 45

Gly Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe  
50 55 60

Gln Gly Arg Val Thr Met Thr Glu Asp Thr Ser Thr Asp Thr Ala Tyr  
65 70 75 80

Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys  
85 90 95

Glu Thr Gly Leu Arg Ser Tyr Asn Tyr Gly Arg Asn Leu Asp Tyr Trp  
100 105 110

Gly Gln Gly Thr Leu Val Thr Val Ser Ser  
115 120

<210> 31

<211> 11

<212> PRT

<213> Homo sapiens

<400> 31

Val Leu Pro Phe Asp Pro Ile Ser Met Asp Val  
1 5 10

<210> 32

<211> 14

<212> PRT

<213> Homo sapiens

<400> 32

Ala Leu Gly Ser Trp Gly Gly Trp Asp His Tyr Met Asp Val  
1 5 10

<210> 33

<211> 5

<212> PRT

<213> Homo sapiens

<400> 33

Gly Tyr Ser Trp Arg  
1 5

<210> 34

<211> 5

<212> PRT

<213> Homo sapiens

<400> 34

Ser Tyr Ala Met His  
1 5

<210> 35

<211> 16

<212> PRT

<213> Homo sapiens

<400> 35

Asp Ile Ser Tyr Ser Gly Ser Thr Lys Tyr Lys Pro Ser Leu Arg Ser  
1 5 10 15

<210> 36

<211> 17

<212> PRT

<213> Homo sapiens

<400> 36

Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 37

<211> 11

<212> PRT

<213> Homo sapiens

<400> 37

Ala Thr Trp Asp Asp Gly Leu Asn Gly Pro Val  
1 5 10

<210> 38

<211> 11

<212> PRT

<213> Homo sapiens

<400> 38

Ala Ala Trp Asp Asp Ser Leu Asn Gly Trp Val  
1 5 10

<210> 39

<211> 13

<212> PRT

<213> Homo sapiens

<400> 39

Ser Gly Ser Ser Ser Asn Ile Arg Ser Asn Pro Val Ser  
1 5 10

<210> 40

<211> 13

<212> PRT

<213> Homo sapiens

<400> 40

Ser Gly Ser Ser Ser Asn Ile Gly Ser Asn Thr Val Asn  
1 5 10

<210> 41

<211> 7

<212> PRT

<213> Homo sapiens

<400> 41

Gly Ser His Gln Arg Pro Ser  
1 5

<210> 42

<211> 7

<212> PRT

<213> Homo sapiens

<400> 42

Ser Asn Asn Gln Arg Pro Ser  
1 5

<210> 43

<211> 16

<212> PRT

<213> Homo sapiens

<400> 43

Val Arg Asp Leu Gly Tyr Arg Val Leu Ser Thr Phe Thr Phe Asp Ile  
1 5 10 15

<210> 44  
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<212> PRT  
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<400> 44  
Asp Gly Arg Ser Gly Ser Tyr Ala Arg Phe Asp Gly Met Asp Val  
1 5 10 15  
Met Gly Ser Ser Val Val Ala Thr Tyr Asn Ala Phe Asp Ile  
1 5 10

<210> 45  
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<400> 45  
Met Gly Ser Ser Val Val Ala Thr Tyr Asn Ala Phe Asp Ile  
1 5 10

<210> 46  
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<400> 46  
Asp Ala Asp Gly Asp Gly Phe Ser Pro Tyr Tyr Phe Pro Tyr  
1 5 10

<210> 47  
<211> 12  
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Leu Arg Asn Asp Gly Trp Asn Asp Gly Phe Asp Ile  
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<210> 48  
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<212> PRT

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<400> 48

Asp Ser Glu Thr Ala Ile Ala Ala Ala Gly Arg Phe Asp Ile  
1 5 10

<210> 49

<211> 13

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<213> Homo sapiens

<400> 49

Glu Asp Gly Thr Thr Val Pro Ser Gln Pro Leu Glu Phe  
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<210> 50

<211> 12

<212> PRT

<213> Homo sapiens

<400> 50

Gly Ser Gly Ser Tyr Leu Gly Tyr Tyr Phe Asp Tyr  
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<210> 51

<211> 13

<212> PRT

<213> Homo sapiens

<400> 51

Gly Leu Arg Ser Tyr Asn Tyr Gly Arg Asn Leu Asp Tyr  
1 5 10

<210> 52

<211> 9

<212> PRT

<213> Homo sapiens

<400> 52

Cys Ser Tyr Val His Ser Ser Thr Asn  
1 5

<210> 53

<211> 9

<212> PRT

<213> Homo sapiens

<400> 53

Gln Val Trp Asp Asn Thr Asn Asp Gln  
1 5

<210> 54

<211> 5

<212> PRT

<213> Homo sapiens

<400> 54

Asn Phe Ala Met Ser  
1 5

<210> 55

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<213> Homo sapiens

<400> 55

Ser Tyr Thr Met His  
1 5

<210> 56

<211> 5

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<213> Homo sapiens

<400> 56

Asp Tyr Ala Leu His  
1 5

<210> 57

<211> 5

<212> PRT

<213> Homo sapiens

<400> 57

Ser His Tyr Trp Ser  
1 5

<210> 58

<211> 5

<212> PRT

<213> Homo sapiens

<400> 58

Thr Tyr Tyr Trp Ser  
1 5

<210> 59

<211> 5

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<213> Homo sapiens

<400> 59

Asp Tyr Gly Met His  
1 5

<210> 60

<211> 5

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<213> Homo sapiens

<400> 60

Ser His Thr Ile Ser  
1 5

<210> 61

<211> 5

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<213> Homo sapiens

<400> 61

Lys Tyr Ala Ile His  
1 5

<210> 62

<211> 5

<212> PRT

<213> Homo sapiens

<400> 62

Glu Leu Ser Met His  
1 5

<210> 63

<211> 17

<212> PRT

<213> Homo sapiens

<400> 63

Gly Ile Ser Gly Gly Leu Leu Thr His Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 64  
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<400> 64  
Gly Ile Ser Gly Gly Leu Leu Thr His Tyr Ala Asn Ser Val Lys  
1 5 10 15

Gly

<210> 65  
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<400> 65  
Gly Ile Thr Pro Ile Phe Gly Thr Val Asn Tyr Ala Gln Lys Phe Gln  
1 5 10 15

Gly

<210> 66  
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<400> 66  
Thr Gly Thr Ser Ser Ala Ile Gly Asn Tyr Asn Phe Val Pro  
1 5 10

<210> 67  
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<400> 67

Gly Gly Tyr Lys Ile Gly Ser Lys Ser Val His  
1 5 10

<210> 68

<211> 7

<212> PRT

<213> Homo sapiens

<400> 68

Glu Gly Ser Lys Arg Pro Ser  
1 5

<210> 69

<211> 7

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<213> Homo sapiens

<400> 69

Glu Asp Ser Tyr Arg Pro Ser  
1 5

<210> 70

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<400> 70

Gly Ile Ser Trp Asp Ser Thr Ser Ile Gly Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 71

<211> 16

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<213> Homo sapiens

<400> 71

Phe Ile Tyr Asp Gly Ala Arg Thr Arg Phe Asn Pro Ser Leu Arg Ser  
10 15  
1 5

<210> 72

<211> 17

<212> PRT

<213> Homo sapiens

<400> 72

Ala Ile Ser Ser Asn Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val Lys  
10 15  
1 5

Gly

<210> 73

<211> 17

<212> PRT

<213> Homo sapiens

<400> 73

Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe Gln  
10 15  
1 5

Gly

<210> 74

<211> 17

<212> PRT

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Gly Ile Ser Gly Gly Leu Leu Thr His Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 75

<211> 30

<212> PRT

<213> Homo sapiens

<400> 75

Arg Phe Thr Ile Ser Arg Asn Asn Ser Arg Asn Thr Val Tyr Leu Gln  
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Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys  
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<213> Homo sapiens

<400> 76

Ala Ile Ser Gly Ser Gly Ser Thr Tyr Tyr Ala Asn Ser Val Lys  
1 5 10 15

Gly

<210> 77

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<400> 77

Leu Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 78

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<400> 78

Gly Ile Ser Trp Asp Ser Thr Ser Ile Gly Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 79

<211> 16

<212> PRT

<213> Homo sapiens

<400> 79

Phe Ile Tyr Asp Gly Ala Arg Thr Arg Phe Asn Pro Ser Leu Arg Ser  
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<400> 80

Thr Gly Thr Ser Ser Ala Ile Gly Asn Tyr Asn Phe Val Pro  
1 5 10

<210> 81

<211> 7

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<400> 81

Glu Gly Ser Lys Arg Pro Ser  
1 5

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<400> 82

Cys Ser Tyr Val His Ser Ser Thr Asn  
1 5

<210> 83

<211> 5

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<213> Homo sapiens

<400> 83

Asp Tyr Gly Met His  
1 5

<210> 84

<211> 17

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<213> Homo sapiens

<400> 84

Ala Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 85

<211> 14

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<213> Homo sapiens

<400> 85

Asp Ser Glu Thr Ala Ile Ala Ala Gly Arg Phe Asp Ile  
1 5 10

<210> 86

<211> 5

<212> PRT

<213> Homo sapiens

<400> 86

Ser His Thr Ile Ser  
1 5

<210> 87

<211> 17

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<213> Homo sapiens

<400> 87

Gly Ile Thr Pro Ile Phe Gly Thr Val Asn Tyr Ala Gln Lys Phe Gln  
1 5 10 15

Gly

<210> 88

<211> 17

<212> PRT

<213> Homo sapiens

<400> 88

Ala Ile Ser Ser Asn Gly Gly Asn Thr Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 89

<211> 12

<212> PRT

<213> Homo sapiens

<400> 89

Gly Ser Gly Ser Tyr Leu Gly Tyr Tyr Phe Asp Tyr  
1 5 10

<210> 90

<211> 13

<212> PRT

<213> Homo sapiens

<400> 90

Gly Leu Arg Ser Tyr Asn Tyr Gly Arg Asn Leu Asp Tyr  
1 5 10

<210> 91

<211> 5

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<213> Homo sapiens

<400> 91  
Ser Tyr Ala Met His  
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<400> 92

Ser Tyr Ala Ile Ser

1 5

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Ser Tyr Gly Met His  
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<400> 94

Glu Leu Ser Met His  
1 5

<210> 95

<211> 17

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<400> 95

Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 96

<211> 17

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<400> 96

Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe Gln  
1 5 10 15

Gly

<210> 97

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<400> 97

Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
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Gly

<210> 98

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Gly Phe Asp Pro Glu Asp Gly Glu Thr Ile Tyr Ala Gln Lys Phe Gln  
1 5 10 15

Gly

<210> 99

<211> 9

<212> PRT

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<400> 99

Gln Val Trp Asp Asn Thr Asn Asp Gln  
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<400> 100  
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1 5 10

<210> 101  
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<400> 101  
Tyr Asp Ser Asp Arg Pro Ser  
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<210> 102  
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<212> PRT  
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<400> 102  
Gln Val Trp Asp Ser Ser Ser Asp His  
1 5

<210> 103  
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<400> 103  
Ser Tyr Ala Met Ser  
1 5

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Ser Tyr Gly Met His  
1 5

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Asp Tyr Ala Met His  
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<400> 106  
Ser Tyr Tyr Trp Ser  
1 5

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<400> 107  
Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 108

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<213> Homo sapiens

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val ile ser Tyr Asp Gly ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

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Gly

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Tyr ile Tyr Tyr Ser Gly Ser Thr Asn Tyr Asn Pro Ser Leu Lys Ser  
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<213> Homo sapiens

<400> 111

Leu Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 112

<211> 16

<212> PRT

<213> Homo sapiens

<400> 112

Tyr Ile Tyr Tyr Ser Gly Asn Thr Asn Tyr Asn Pro Ser Leu Lys Ser  
1 5 10 15

<210> 113

<211> 17

<212> PRT

<213> Homo sapiens

<400> 113

Ala Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 114

<211> 17

<212> PRT

<213> Homo sapiens

<400> 114

Gly Ile Ser Trp Asp Ser Gly Thr Ile Gly Tyr Ala Asp Ser Val Lys  
1 5 10 15

Gly

<210> 115

<211> 11

<212> PRT

<213> Homo sapiens

<400> 115

Gly Gly Tyr Lys Ile Gly Ser Lys Ser Val His  
1 5 10

<210> 116

<211> 7

<212> PRT

<213> Homo sapiens

<400> 116

Glu Asp Ser Tyr Arg Pro Ser  
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<210> 117

<211> 18

<212> DNA

<213> Artificial sequence

<220>

<223> Description of Artificial sequence: Synthetic Oligonucleotide

<400> 117  
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18

<210> 118

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Oligonucleotide

<400> 118

ggccgcaaat tctatttcaa gg

22

<210> 119

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Oligonucleotide

<400> 119

gagacacacc agtgtggc

18

<210> 120

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic Oligonucleotide

<400> 120

cacaacagag gcagttcc

18

<210> 121

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic Oligonucleotide

<400> 121  
ctaaactagc tagtctcc

18

<210> 122

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<212> PRT

<213> Homo sapiens

<400> 122

Gly Gly Tyr Lys Ile Gly Ser Lys Ser Val His  
1 5 10

<210> 123

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<212> PRT

<213> Homo sapiens

<400> 123

Gln Val Lys Leu Leu Glu Ser Gly Gly Leu Val Gln Pro Gly Arg  
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Asp Asp Tyr  
20 25 30

Ala Leu His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val  
35 40 45

Ser Gly Ile Ser Trp Asp Ser Thr Ser Ile Gly Tyr Ala Asp Ser Val  
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr  
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Leu Tyr Tyr Cys  
85 90 95

Val Lys Asp Met Gly Ser Ser Val Val Ala Thr Tyr Asn Ala Phe Asp  
100 105 110

Ile Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser  
115 120

<210> 124  
<211> 14  
<212> PRT  
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<400> 124  
Thr Gly Thr Ser Ser Asp Val Gly Ser Tyr Asn Leu Val Ser  
1 5 10

<210> 125  
<211> 7  
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<213> Homo sapiens

<400> 125  
Glu Val Ser Lys Arg Pro Ser  
1 5

<210> 126  
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<212> PRT  
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<400> 126  
Cys Ser Tyr Ala Gly Ser Ser Thr Phe  
1 5

<210> 127  
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<212> PRT  
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<400> 127  
Lys Tyr Ala Ile His  
1 5

<210> 128

<211> 7

<212> PRT

<213> *Homo sapiens*

<400> 128

Arg Asn Asn Gln Arg Pro Ser  
1 5